

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT:
- (A) NAME: Transgene S.A.
 - (B) STREET: 11 rue de Molsheim
 - (C) CITY: Strasbourg
 - (E) COUNTRY: France
 - (F) POSTAL CODE: 67082
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- (ii) TITLE OF INVENTION: New method for the preparation of a viral vector
- (iii) NUMBER OF SEQUENCES: 10
- (iv) COMPUTER READABLE FORM:
- (A) MEDIUM TYPE: Tape
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)

(2) INFORMATION FOR SEQ ID NO: 1:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 38 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iii) ANTI-SENSE: NO
- (vi) ORIGINAL SOURCE:
- (A) ORGANISM: human adenovirus
 - (B) STRAIN: serotype 5
 - (C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG6597
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

GCCGAATTCT TAATTAACAT CATCAATAAT ATACCTTA

(2) INFORMATION FOR SEQ ID NO: 2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 35 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iii) ANTI-SENSE: YES
- (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: human adenovirus
 - (B) STRAIN: serotype 5
 - (C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG6598
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

GACAGATCTG TCGACGTGGC AGGTAAGATC GATCA

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(2) INFORMATION FOR SEQ ID NO: 3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 35 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iii) ANTI-SENSE: NO
- (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: human adenovirus
 - (B) STRAIN: serotype 5
 - (C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG6599
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

AGGAGATCTG TCGACTCTCA AACATGTCTG CGGGT

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(2) INFORMATION FOR SEQ ID NO: 4:

09338491-082701

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 47 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iii) ANTI-SENSE: NO
- (vi) ORIGINAL SOURCE:
 (A) ORGANISM: Rous sarcoma virus
 (C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG5892
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

GTCGTAGGAT CCAGCTGCTC CCTGCTTGTG TGTTGGAGGT CGCTGAG 47

(2) INFORMATION FOR SEQ ID NO: 5:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 47 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iii) ANTI-SENSE: YES
- (vi) ORIGINAL SOURCE:
 (A) ORGANISM: Rous sarcoma virus
 (C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG5893
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

GTCAGTGACG TCCAGGTGTC ACACCAATGT GGTGAATGGT CAAATGG 47

(2) INFORMATION FOR SEQ ID NO: 6:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 25 base pairs
 (B) TYPE: nucleic acid

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- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iii) ANTI-SENSE: NO

(vi) ORIGINAL SOURCE:

- (A) ORGANISM: human adenovirus
- (B) STRAIN: serotype 5
- (C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG5455

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

ATCGGAATTC AAGATGATTA GGTAC

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(2) INFORMATION FOR SEQ ID NO: 7:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iii) ANTI-SENSE: YES

(vi) ORIGINAL SOURCE:

- (A) ORGANISM: human adenovirus
- (B) STRAIN: serotype 5
- (C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG5456

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

ATCGTCTAGA TTAAGGCATT TTCTTTTC

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(2) INFORMATION FOR SEQ ID NO: 8:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 36 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single

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(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iii) ANTI-SENSE: NO

(vi) ORIGINAL SOURCE:

(A) ORGANISM: canine adenovirus

(B) STRAIN: CAV-2

(C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG6974

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

CAGGGATCCG CGGCCGCATC ATCAATAATA TACAGG

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(2) INFORMATION FOR SEQ ID NO: 9:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 29 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iii) ANTI-SENSE: YES

(vi) ORIGINAL SOURCE:

(A) ORGANISM: canine adenovirus

(B) STRAIN: CAV-2

(C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG6975

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

CTGCTGCAGT CAGAAATGCT AGCAGGAGA

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(2) INFORMATION FOR SEQ ID NO: 10:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 27 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

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(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iii) ANTI-SENSE: NO

(vi) ORIGINAL SOURCE:

(A) ORGANISM: canine adenovirus

(B) STRAIN: CAV-2

(C) INDIVIDUAL ISOLATE: synthetic oligonucleotide OTG6976

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

TGCGGATCCA CAGACTAAGC GGAGGTA

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